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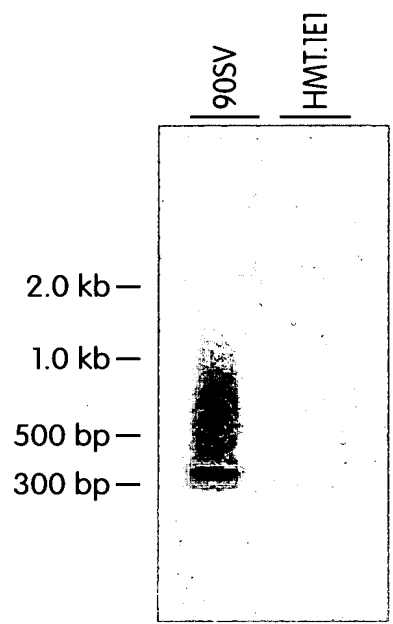


Fig. 1A

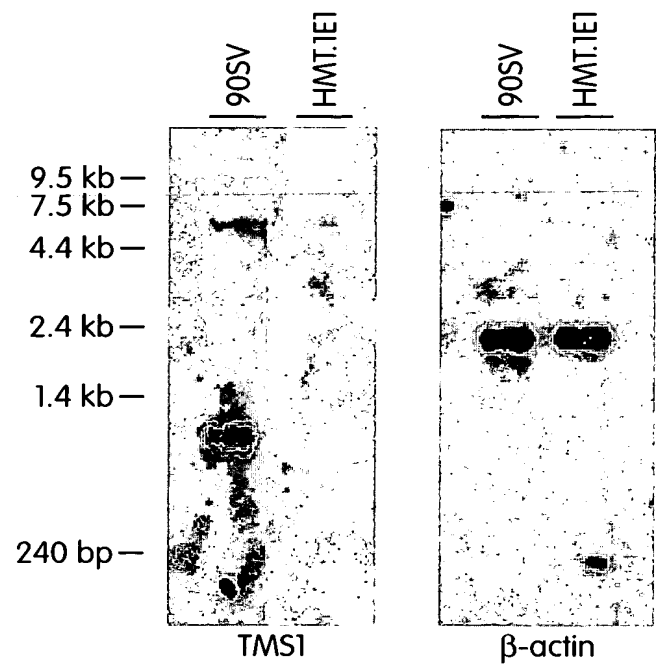


Fig. 1B

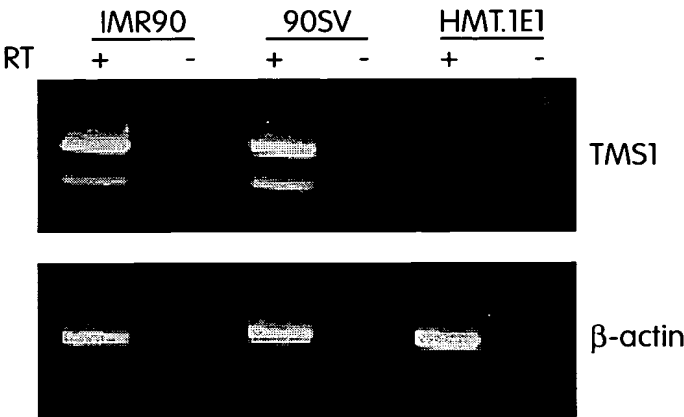


Fig. 1C

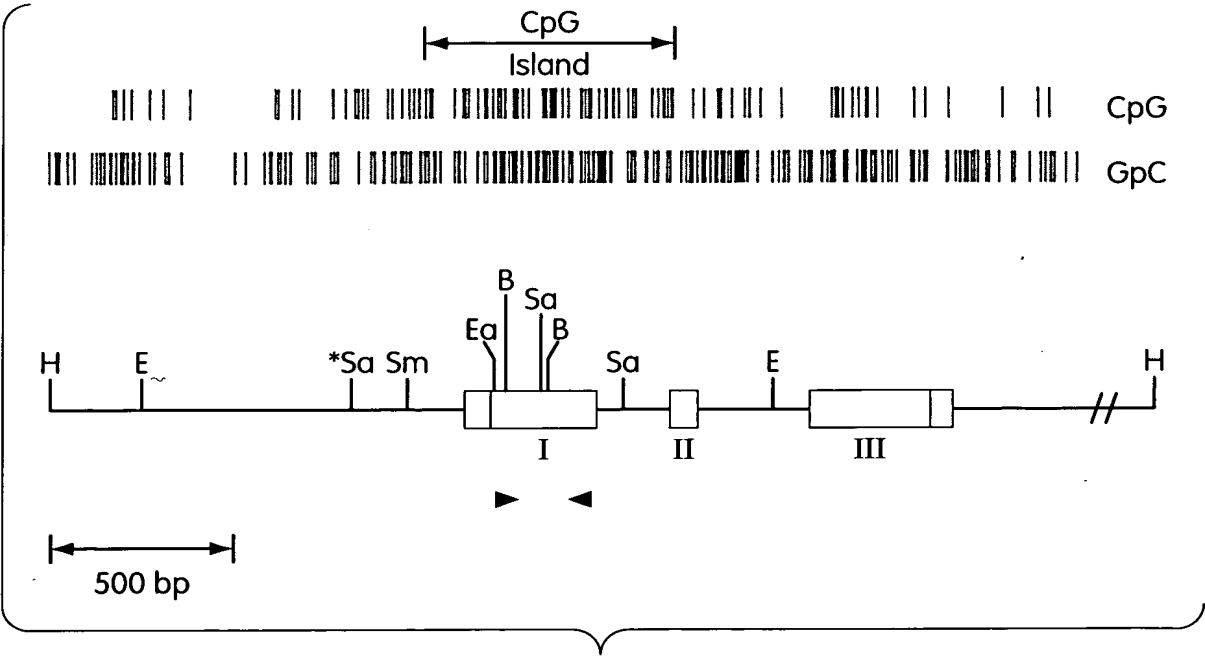
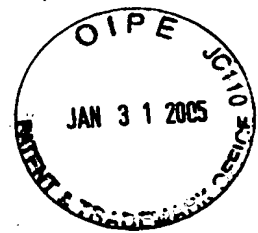


Fig. 2A



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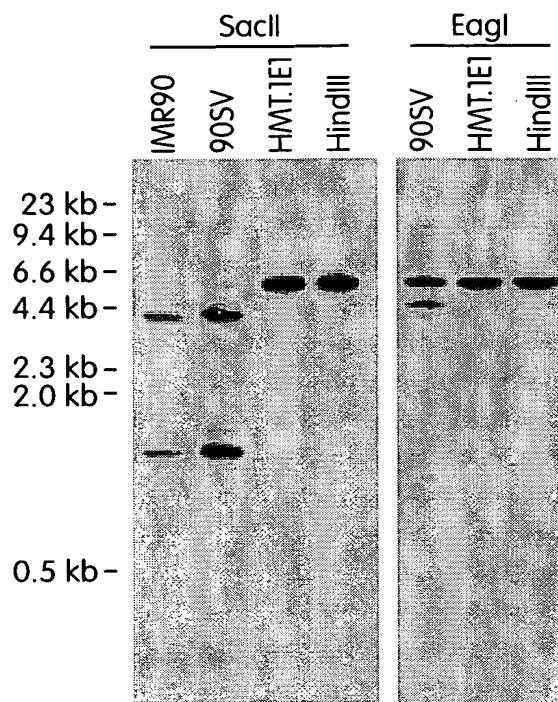


Fig. 2B

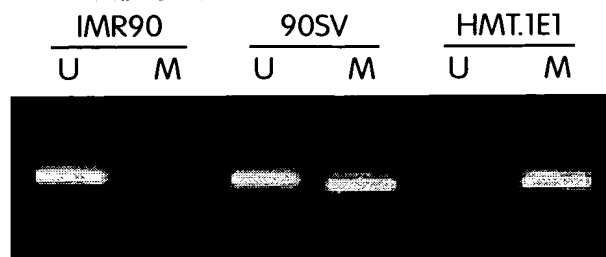


Fig. 2C

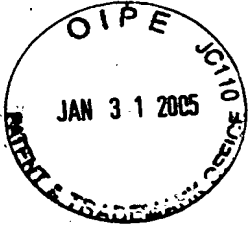


Fig. 3A

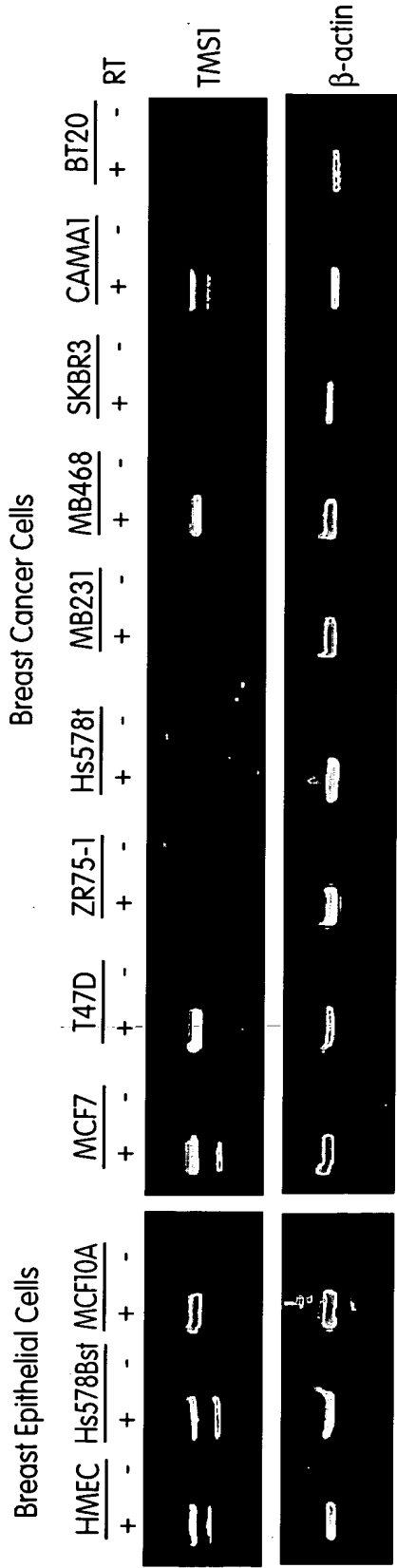
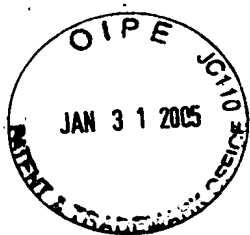


Fig. 3B

TMS1

β-actin



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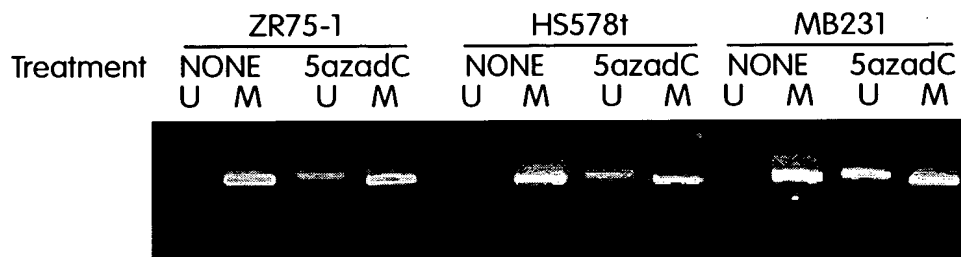


Fig. 3C

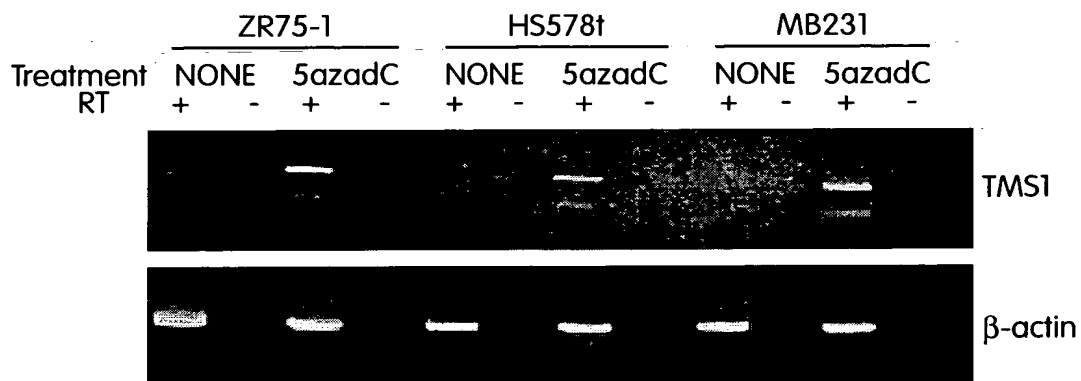
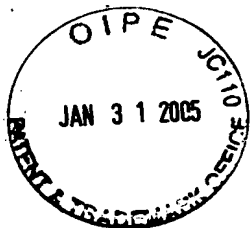


Fig. 3D



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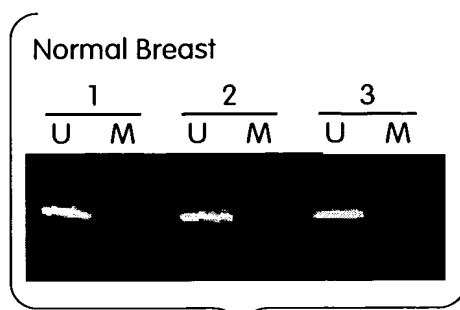


Fig. 4A

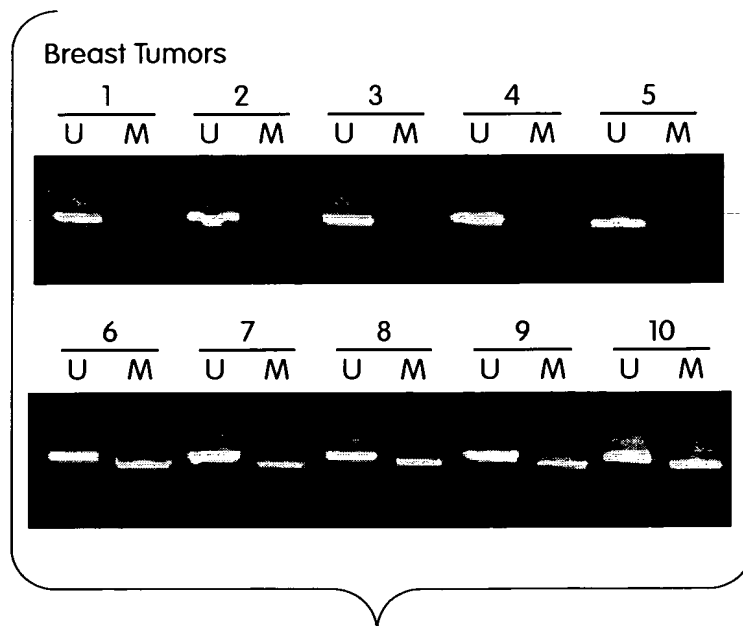
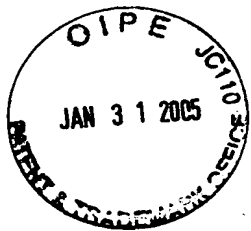


Fig. 4B



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Tumor/Normal Pairs

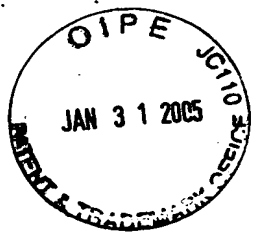
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U	M	U	M	U	M	U	M	U	M	U	M



7N	7T	8N	8T	9N	9T	10N	10T	11N	11T	12N	12T
U	M	U	M	U	M	U	M	U	M	U	M



Fig. 4C



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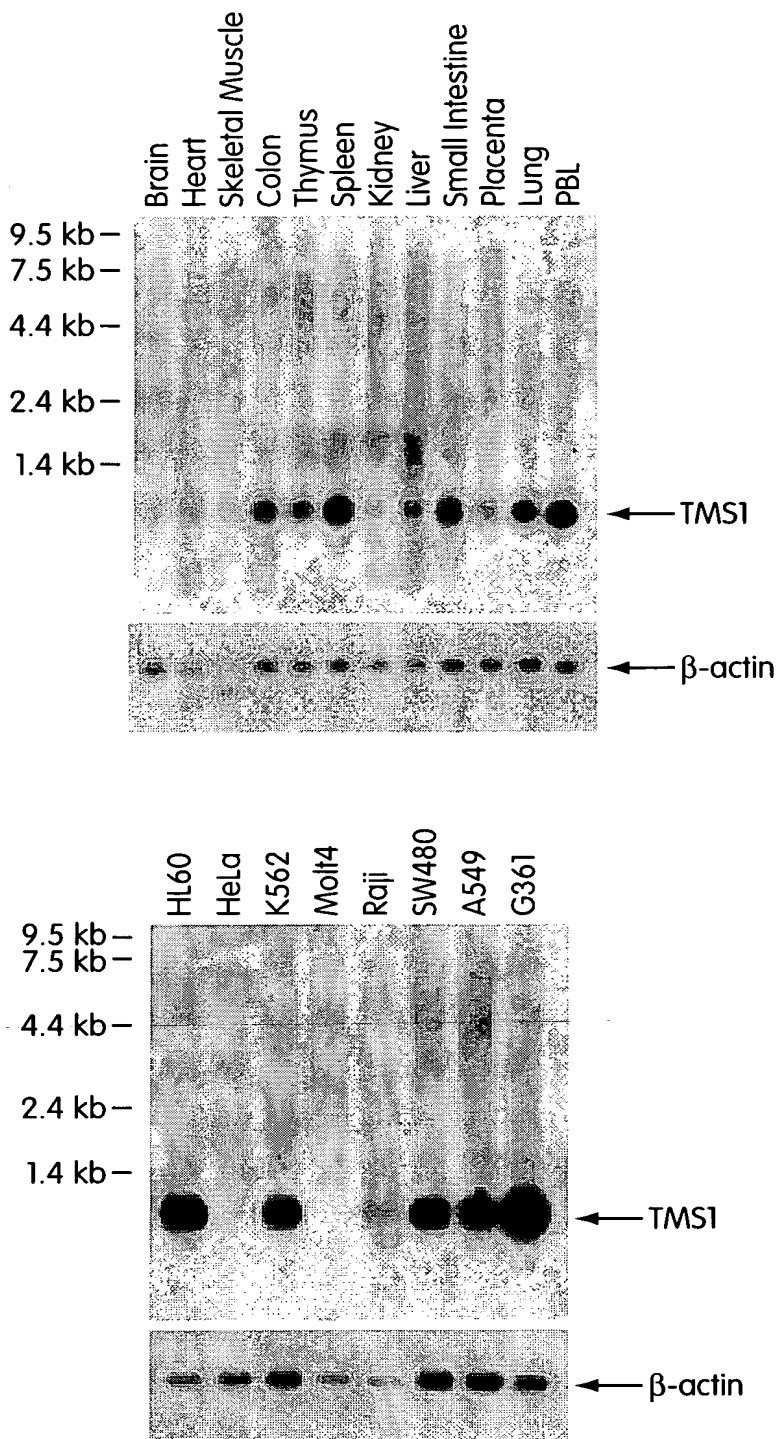
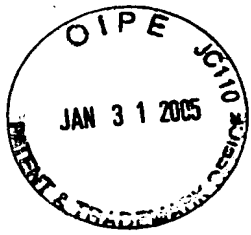


Fig. 5A



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AAKPGHLHFDQRAALARVTN--VEMEDDAFYG-KVETDEQYQAVRAE--PAPPSKMKKIFSTPANNWTCCKLLQAR--ESQSYLVEDZERS
ESHPHIQLEKSNRELVTHIRN--TQCLAVNLEKNDYFSAEDAEIVCAC--PQPDKVRKIIFDLVQSKCEEVSEFEYLLQQADAYVDLRPWLL
QPGIAQQWQSKREDIVNQTEACLNSIDALSRDLIMREDYELVSTK--PRTSKVNOFHTTDIQCEEFAKIVQKFK--DNKQMGLOPYPEI
---MADKVKERKLFIRSMGEGTINGHDELLQTRVANKREEMKVKREN-ALVMDTTRALHDSVLPKCAQCOQCTTYTC--EEDSYLAGTGLS
MHPHQETKKNVWFAKQLL--SESEHEHEKDLZTLEMRLLQAK--VGSFSQNVETLILPKRCPQAFDAACEAR--ETKQGHLEDMLLT
DEADRLRRCRURVVEELQV--DQLEWNVHLSRELFRPHMIEDIQRAGSGSRDDAQOQZIDLETRECSQALPLRISCE--DTGQDMIASFLRT
IASDDISLZIKNMAFQQLTC--VLPFDNLKANKVZINKQEHDIKQK--TQIPLOARETDTILVKNAANIKKNCFK--EIDSTLYKNLFFVQ
MEARDKQVRSIRUFGAEVLV--EGLMIOYLYQEGLETENHIOELNAQ--TIGLRKTMLEHDLPSRCPKFDTEHDSIQ--EFPWVREKLKAR
LFEVKDAEENLRVYCEKELIA--ERHFDHURAKKLSREDEEESCR--HSSRKFRAGKGGVYIQ--ENPKGLDTLVESIRREKTONFYIQKITD
WDAKARNCELOHFEAEFEKDIKT--SYTMDHMTSDGFTTISEPEKVRNE--PQOQQAAMFKMLKNDNSIVSSEYNALEH--GYKDLAALZHDG

TMS1

CARD4

RICK

procaspase-1

procaspase-2

procaspase-9

CIAP-1

RAIDD

Bc110

Apaf-1

Fig. 5B



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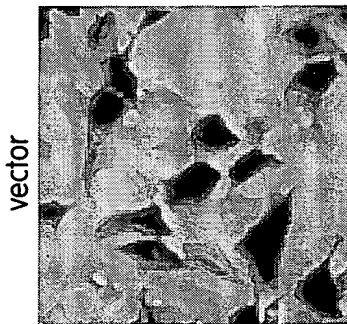
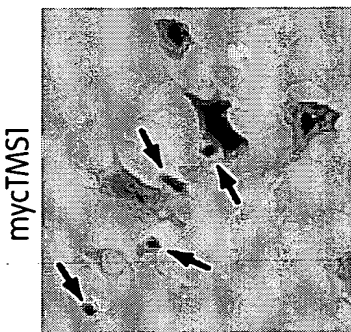
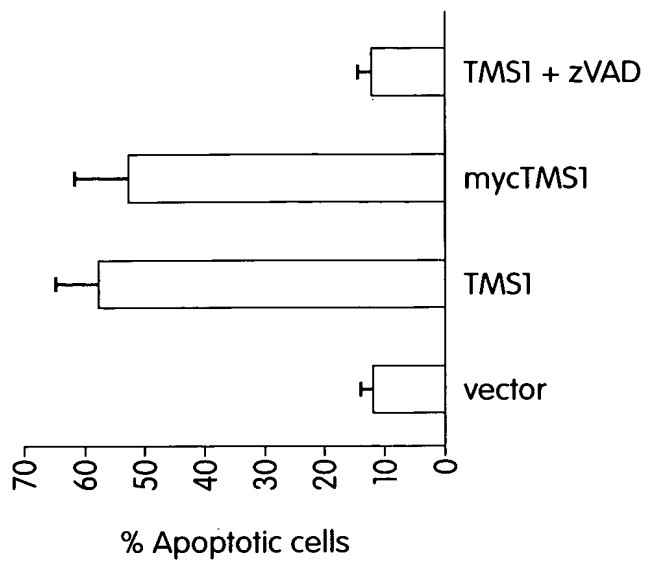
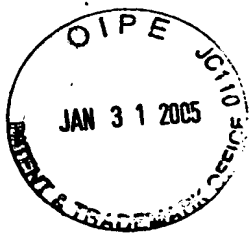


Fig. 5C



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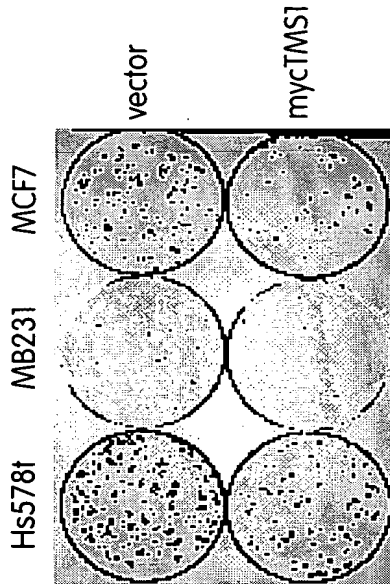
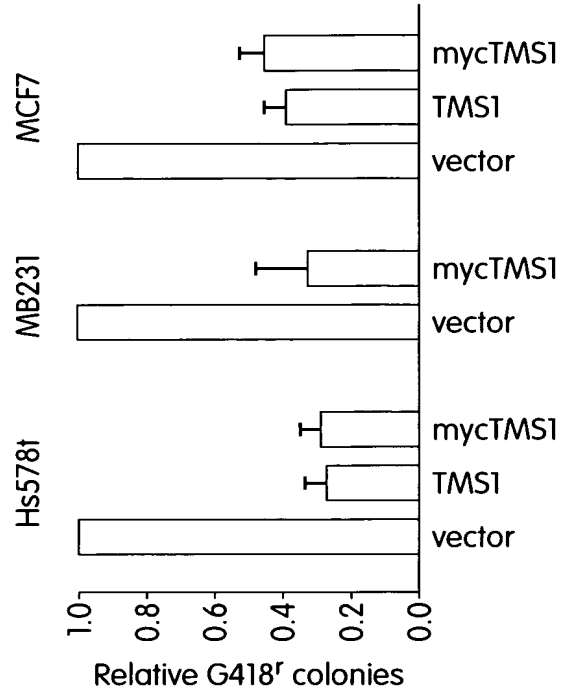
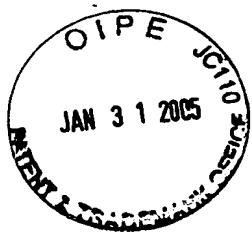


Fig. 5D



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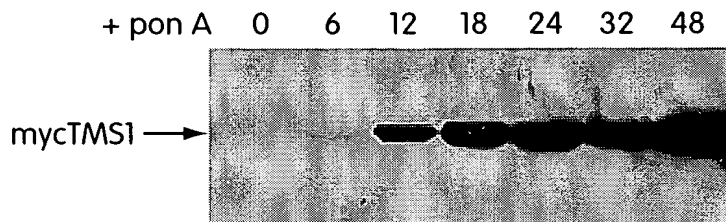


Fig. 6A

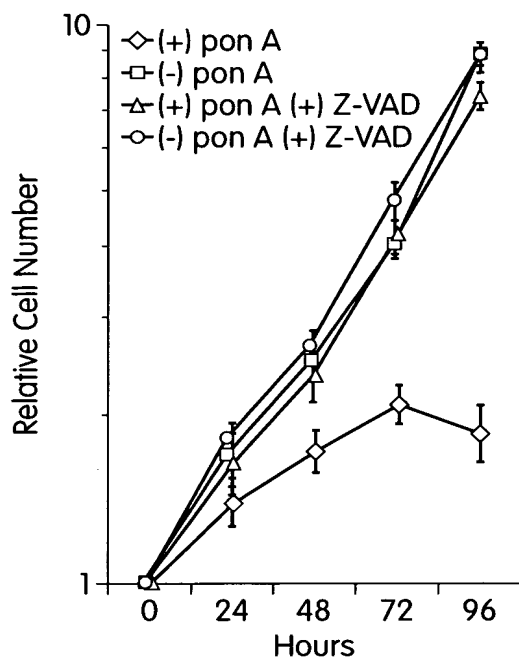


Fig. 6B

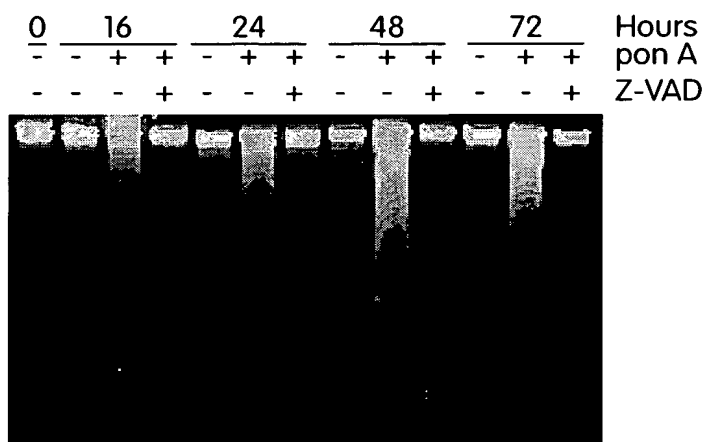
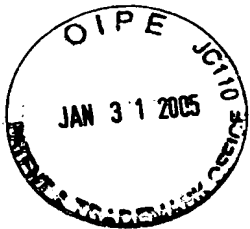


Fig. 6C



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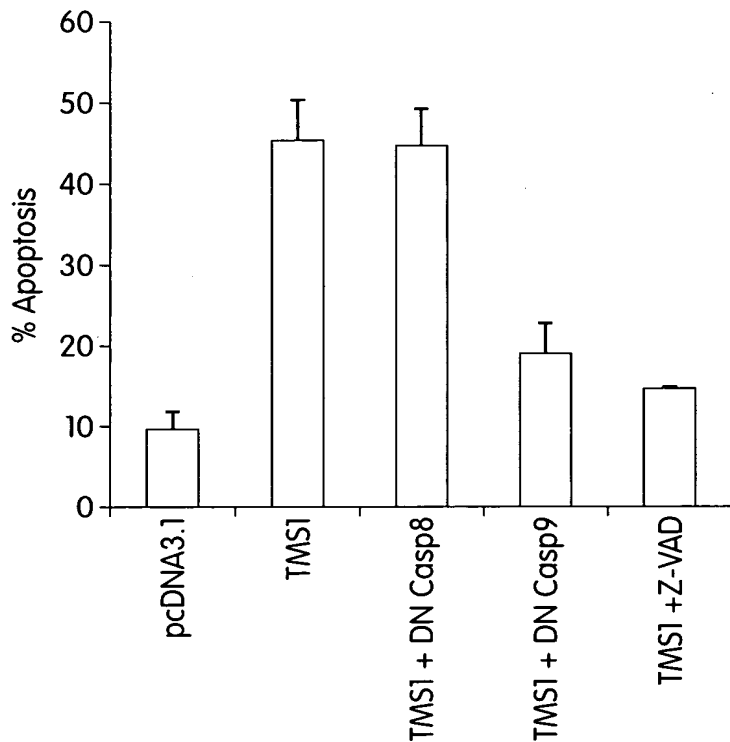


Fig. 7A

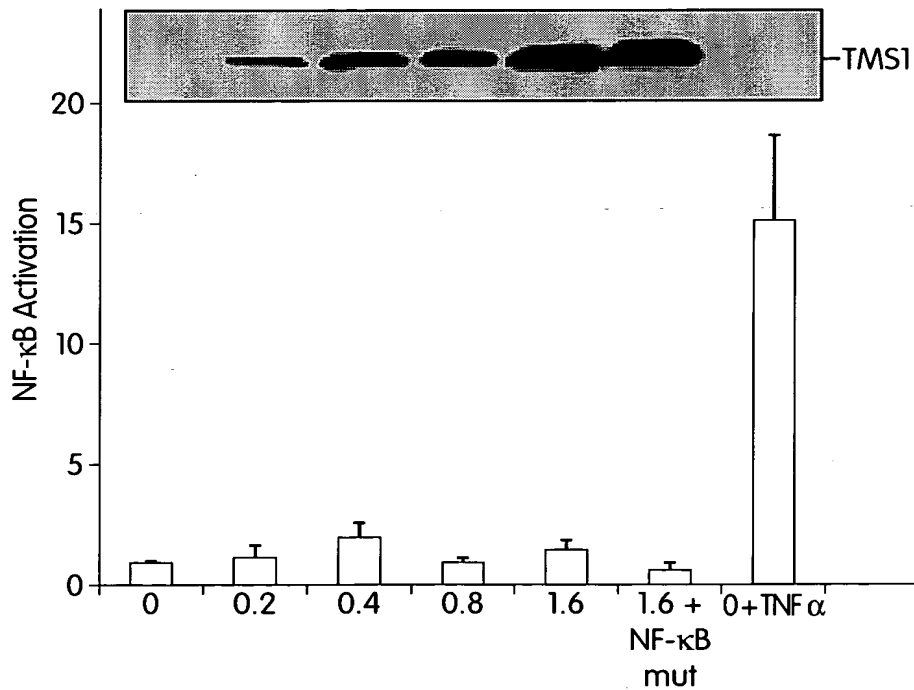


Fig. 7B

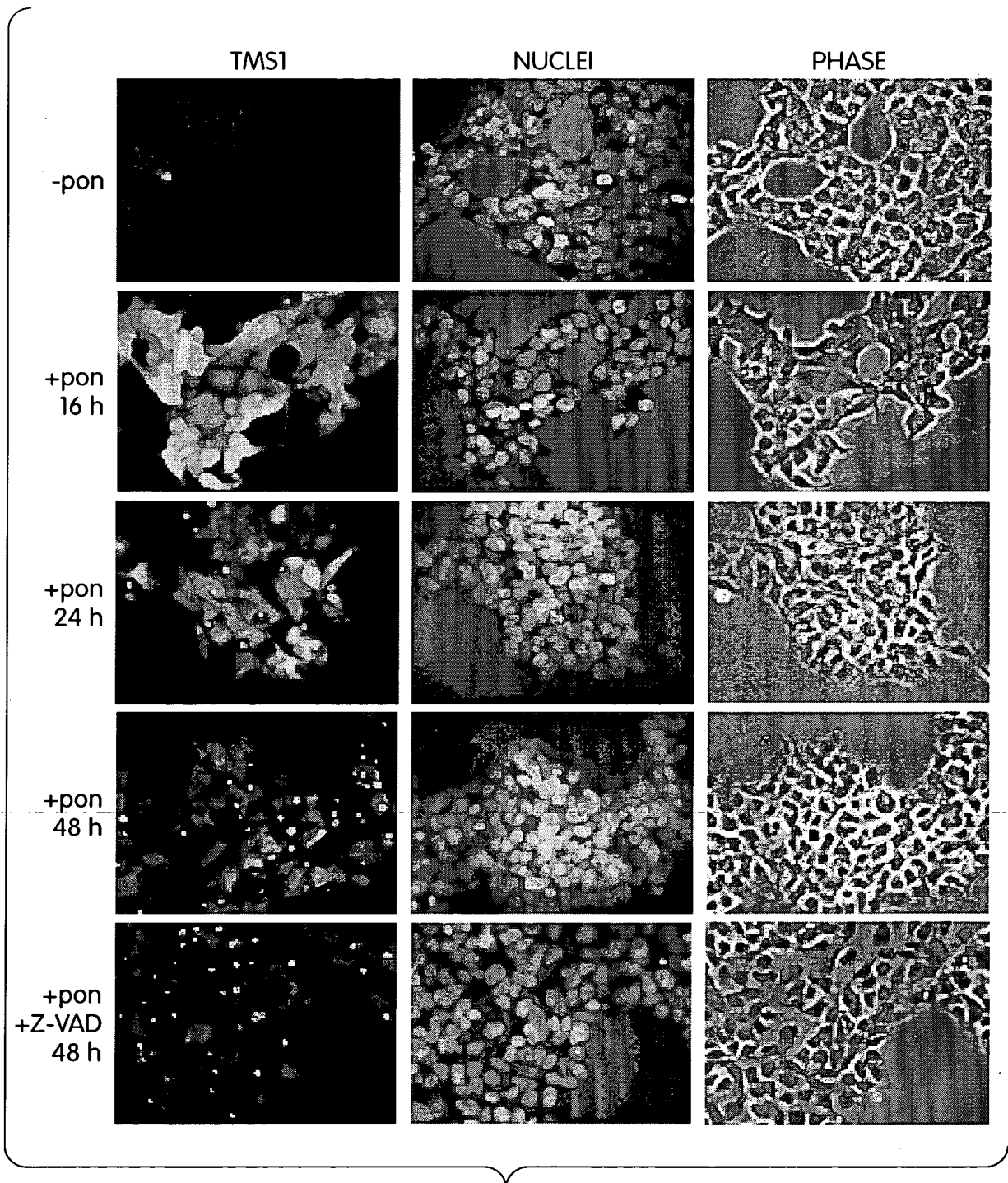
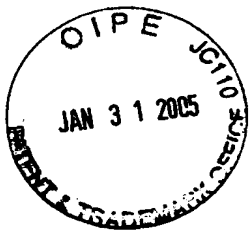


Fig. 8A

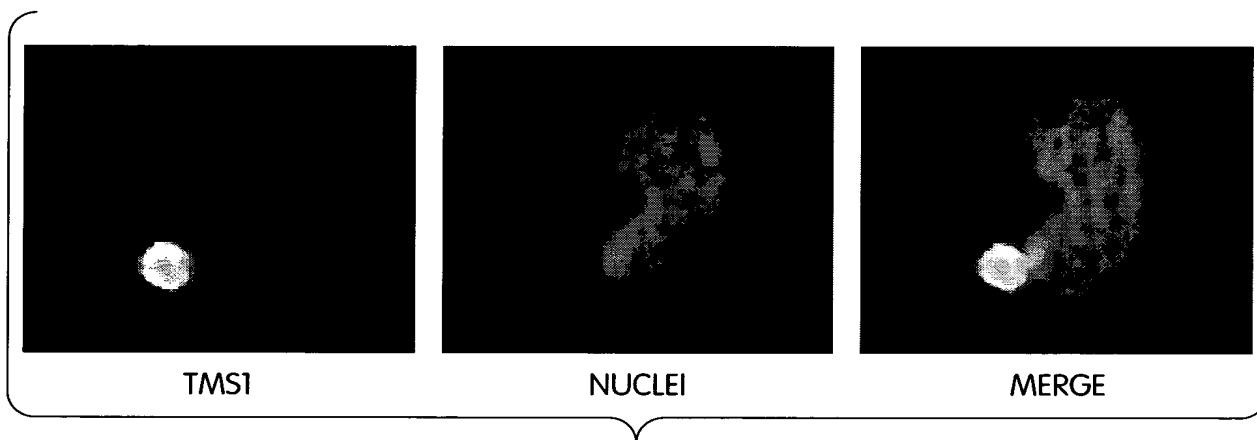
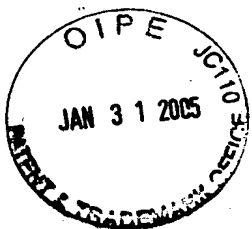
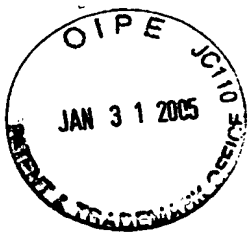


Fig. 8B



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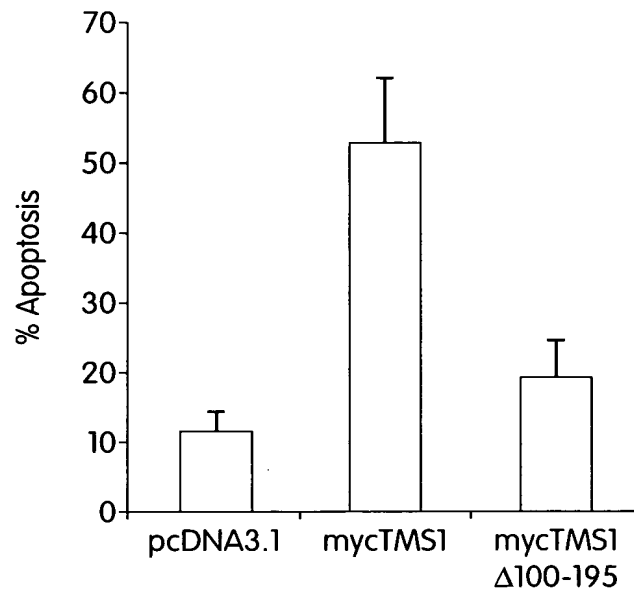


Fig. 9A

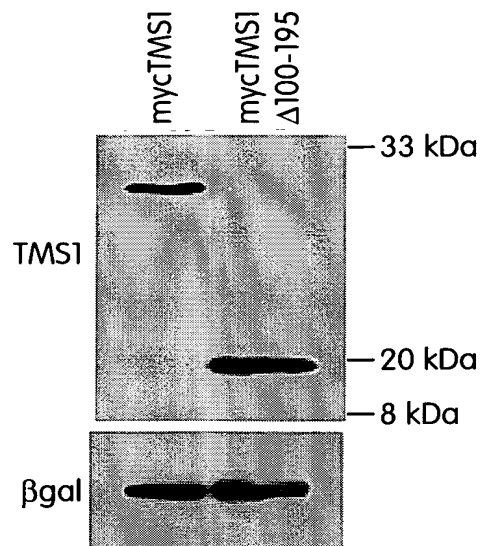
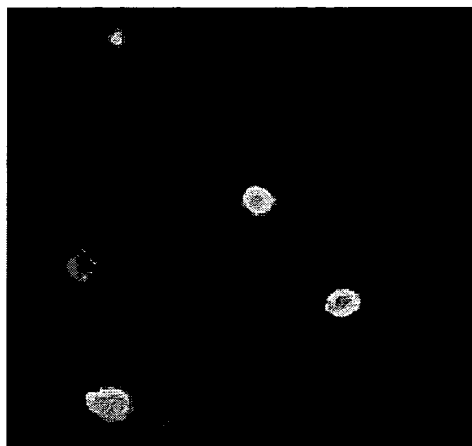


Fig. 9B



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myc TMS1



myc TMS1 Δ 100-195

<u>Construct</u>	<u>Aggregates</u>	<u>Cytoplasmic Staining</u>
mycTMS1	303 (55%)	249 (45%)
mycTMS1 Δ 100-195	7 (1%)	511 (99%)

Fig. 9C

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